

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/896,896
Source: O/P/E
Date Processed by STIC: 2/9/2002

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/896,896

DATE: 02/07/2002

TIME: 13:36:42

Input Set : A:\31-17US Sequence listing.txt

Output Set: N:\CRF3\02072002\I896896.raw

**Does Not Comply
Corrected Diskette Needed**

3 <110> APPLICANT: MAXYGEN APS
5 <120> TITLE OF INVENTION: N-TERMINALLY EXTENDED POLYPEPTIDES
7 <130> FILE REFERENCE: 0217us210
W--> 9 <140> CURRENT APPLICATION NUMBER: US/09/896,896
W--> 0 <160> NUMBER OF SEQ ID NOS:
9 <170> SOFTWARE: PatentIn Ver. 2.1

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ERRORED SEQUENCES

see next page

09/896,896 2

SEQUENCE LISTING

RECEIVED

MAY 20 2002

<110> MAXYGEN APS

<120> N-TERMINALLY EXTENDED POLYPEPTIDES

<130> 0217us210

<170> PatentIn Ver. 2.1

<210> 1

→ <160>

TECH CENTER 1600/2000

this is a mandatory

numeric identifier and

needs a response

FYI: 1/3 sequence in
submitted file

Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to locate a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

FYI

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/896,896

DATE: 02/07/2002

TIME: 13:36:44

Input Set : A:\31-17US Sequence listing.txt

Output Set: N:\CRF3\02072002\I896896.raw

L:9 M:282 W: Numeric Field Identifier Missing, <140> CURRENT APPLICATION NUMBER: is Added.
L:0 M:282 W: Numeric Field Identifier Missing, <160> is required.
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:156 M:283 W: Missing Blank Line separator, <220> field identifier
L:160 M:283 W: Missing Blank Line separator, <400> field identifier
L:392 M:283 W: Missing Blank Line separator, <220> field identifier
L:396 M:283 W: Missing Blank Line separator, <400> field identifier
L:611 M:283 W: Missing Blank Line separator, <400> field identifier
L:636 M:283 W: Missing Blank Line separator, <400> field identifier
L:664 M:283 W: Missing Blank Line separator, <220> field identifier
L:668 M:283 W: Missing Blank Line separator, <400> field identifier
L:915 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:934 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:953 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
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L:2408 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99
L:2446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:101
L:0 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (0) Counted (113)